

FIGURE 1

	1		50
Bcl-2/iso1	MAHAGRTGYD NREIVMKYIH YKLSQRGYEW DAGDVGAAPP GAAPAPGIFS		
Bcl-2/iso2	MAHAGRTGYD NREIVMKYIH YKLSQRGYEW DAGDVGAAPP GAAPAPGIFS		
Bcl-2/iso3	MAHAGRTGYD NREIVMKYIH YKLSQRGYEW DAGDVGAAPP GAAPAPGFFS		
Bcl-xL	MSMAMSQS NRELVVDFLS YKLSQKGYSW SQFSDVEENR TEAPEETESE		
	51		100
Bcl-2/iso1	SQPGHTPHPA ASRDPVARTS PLQTPAAPGA AAGPALSPVP PVVHLALRQA		
Bcl-2/iso2	SQPGHTPHPA ASRDPVARTS PLQTPAAPGA AAGPALSPVP PVVHLTLRQA		
Bcl-2/iso3	SQPGHTPHPA ASRDPVARTS PLQTPAAPGA AAGPALSPVP PVVHLTLRQA		
Bcl-xL	METPSAINGN PSWHLADSPA VNGATGHSSS LDAREVIP-M AAVKQALREA		
	101		150
Bcl-2/iso1	GDDFSRRYRG DFAEMSSQLH LTPFTARGRF ATVVEELFRD GVNWGRIVAF		
Bcl-2/iso2	GDDFSRRYRR DFAEMSSQLH LTPFTARGRF ATVVEELFRD GVNWGRIVAF		
Bcl-2/iso3	GDDFSRRYRR DFAEMSSQLH LTPFTARGRF ATVVEELFRD GVNWGRIVAF		
Bcl-xL	GDEFELRYRR AFSDLTSQLH ITPGTAYQSF EQVVNELFRD GVNWGRIVAF		
	151		200
Bcl-2/iso1	FEFGGVMCVE SVNREMSPLV DNIALWMTEY LNRHLHTWIQ DNGGWDAFVE		
Bcl-2/iso2	FEFGGVMCVE SVNREMSPLV DNIALWMTEY LNRHLHTWIQ DNGGWDAFVE		
Bcl-2/iso3	FEFGGVMCVE SVNREMSPLV DNIALWMTEY LNRHLHTWIQ DNGGWDAFVE		
Bcl-xL	FSFGGALCVE SVDKEMQVLV SRIAAMWATY LNDHLEPWIQ ENGGWDTFVE		
	201		239
Bcl-2/iso1	LYGPSMRPLF DFSWLSLCTL LSLALVGACI TLGAYLGHK		
Bcl-2/iso2	LYGPSMRPLF DFSWLSLCTL LSLALVGACI TLGAYLGHK		
Bcl-2/iso3	LYGPSMRPLF DFSWLSLCTL LSLALVGACI TLGAYLGHK		
Bcl-xL	LYGNNAAAES RKGQERFNRW FLTGMTVAGV VLLGSLFSRK		

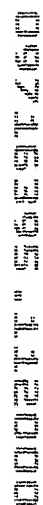
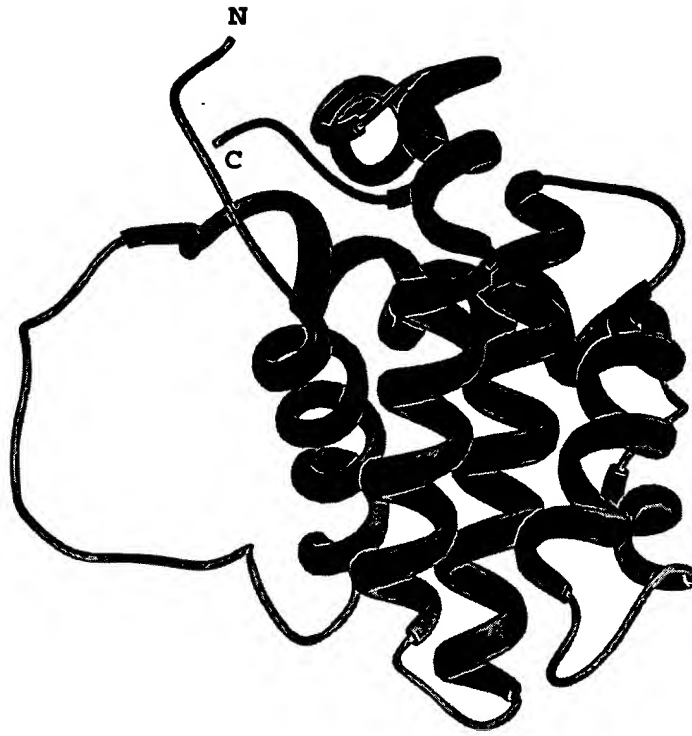
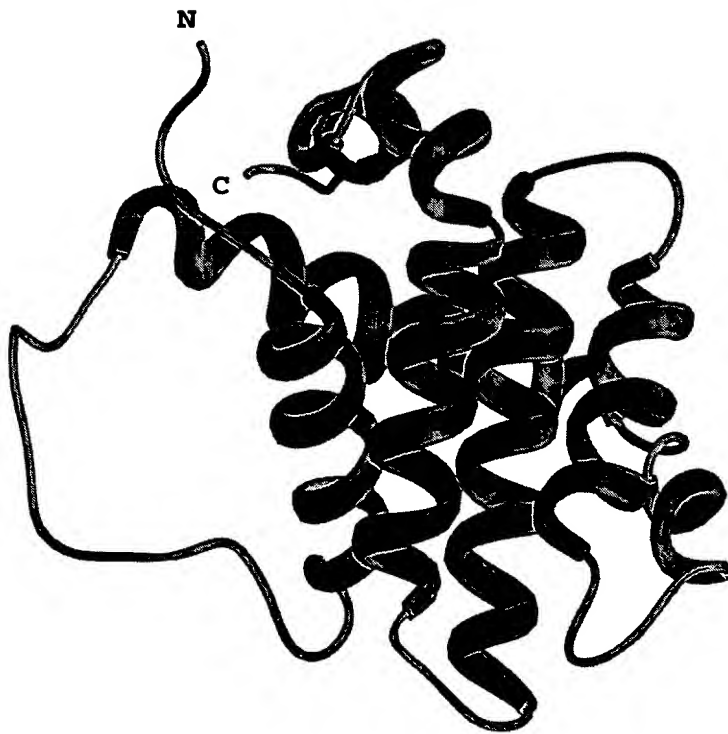


FIGURE 3

A



B



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